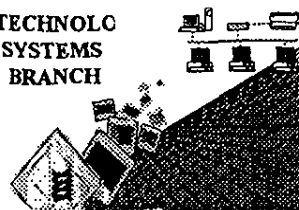


5040

BIOTECHNOLOGY
SYSTEMS
BRANCH



RAW SEQUENCE LISTING
ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/030,658
Source: PGF/10
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FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216.

PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax)

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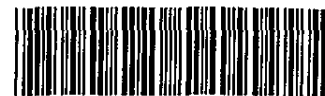
Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

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2011 South Clark Place, Arlington, VA 22202
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Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 01/29/2002



PCT10

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/030,658

DATE: 02/08/2002
TIME: 10:31:52

Input Set : A:\PTO.VSK.txt
Output Set: N:\CRF3\02082002\J030658.raw

Does Not Comply
Corrected Diskette Needed

3 <110> APPLICANT: Yamamura Ken-ichi
4 Araki Kimi
6 <120> TITLE OF INVENTION: TRAPVECTOR
8 <130> FILE REFERENCE: PH-976PCT
10 <140> CURRENT APPLICATION NUMBER: US/10/030,658
10 <141> CURRENT FILING DATE: 2002-01-11
10 <150> PRIOR APPLICATION NUMBER: JP99/200997
11 <151> PRIOR FILING DATE: 1999-07-14
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15 <170> SOFTWARE: PatentIn Ver. 2.0

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128 <211> LENGTH: 400
129 <212> TYPE: DNA
130 <213> ORGANISM: Mus musculus
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/030,658

DATE: 02/08/2002
TIME: 10:31:52

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Output Set: N:\CRF3\02082002\J030658.raw

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Item 9

last sequence in file

delete

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/030,658

DATE: 02/08/2002

TIME: 10:31:53

Input Set : A:\PTO.VSK.txt

Output Set: N:\CRF3\02082002\J030658.raw

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L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date
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M:340 Repeated in SeqNo=11
L:147 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:12
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L:161 M:340 E: (46) "n" or "Xaa" used: Feature required, for SEQ ID#:13
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L:177 M:212 E: (34) Invalid or duplicate Sequence ID Number, SEQUENCE ID NOS:14 differs:12
L:184 M:254 E: No. of Bases conflict, LENGTH:Input:1 Counted:212 SEQ:14
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